



<110> Renauld, Jean-Christophe
Dumoutier, Laure

<120> Isolated Nucleic Acid Molecules Which Encode A Soluble IL-TIF/IL-22
Receptor or Binding Protein Which Binds to IL-TIF/IL-22, And Uses Thereof

<130> LUD 5684.2 (10106926)

<140> US 09/919,162

<141> 2001-31-07

<150> US 60/245,495

<151> 2000-03-11

<150> US 60/234,583

<151> 2000-09-22

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<212> DNA

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actggcaaca	gcagtgtcta	ttttgtgcag	tacaaaatat	atggacagag	acaatggaaa	240
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tttataatta	acaattcact	agaaaaggag	caaaagggtt	atgaaggggc	tcacagagcg	600
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cccatgttag	acagaagaag	tcagagaagt	gaagagagat	gtgtggaaat	tccatgactt	720
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			20					25					30		
Arg	Val	Gln	Phe	Gln	Ser	Arg	Asn	Phe	His	Asn	Ile	Leu	Gln	Trp	Gln
		35					40					45			
Pro	Gly	Arg	Ala	Leu	Thr	Gly	Asn	Ser	Ser	Val	Tyr	Phe	Val	Gln	Tyr
	50					55					60				
Lys	Ile	Tyr	Gly	Gln	Arg	Gln	Trp	Lys	Asn	Lys	Glu	Asp	Cys	Trp	Gly
65				70					75					80	
Thr	Gln	Glu	Leu	Ser	Cys	Asp	Leu	Thr	Ser	Glu	Thr	Ser	Asp	Ile	Gln
			85					90					95		
Glu	Pro	Tyr	Tyr	Gly	Arg	Val	Arg	Ala	Ala	Ser	Ala	Gly	Ser	Tyr	Ser
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Glu	Trp	Ser	Met	Thr	Pro	Arg	Phe	Thr	Pro	Trp	Trp	Glu	Thr	Lys	Ile
	115						120					125			
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	130				135						140				
Ile	Leu	His	Ala	Pro	Asn	Leu	Pro	Tyr	Arg	Tyr	Gln	Lys	Glu	Lys	Asn
145					150					155				160	
Val	Ser	Ile	Glu	Asp	Tyr	Tyr	Glu	Leu	Leu	Tyr	Arg	Val	Phe	Ile	Ile
			165					170					175		
Asn	Asn	Ser	Leu	Glu	Lys	Glu	Gln	Lys	Val	Tyr	Glu	Gly	Ala	His	Arg
		180						185					190		
Ala	Val	Glu	Ile	Glu	Ala	Leu	Thr	Pro	His	Ser	Ser	Tyr	Cys	Val	Val
	195						200					205			
Ala	Glu	Ile	Tyr	Gln	Pro	Met	Leu	Asp	Arg	Arg	Ser	Gln	Arg	Ser	Glu
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 tttcacaaca ttttgcaatg gcagcctggg agggcactta ctggcaacag cagtgtctat 180
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 ggatgctggc agcacatttc ttgtaacttc ccaggctgca gaacattggc taaatatgga 300
 cagagacaat ggaaaaataa agaagactgt tggggacttc aagaactctc ttgtgacctt 360
 accagtgaac cctcagacat acaggaacct tattacggga ggggtgagggc ggccctcggt 420
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 gatcctccag tcatgaatat aacccaagtc aatggctctt tgttggtaat tctccatgct 540
 ccaaattttac catatagata ccaaaaggaa aaaaatgtat ctatagaaga ttactatgaa 600
 ctactatacc gagtttttat aattaacaat tctactagaa aggagcaaaa ggtttatgaa 660
 ggggctcaca gagcggttga aattgaagct ctaacaccac actccagcta ctgtgtagt 720
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 20 25 30
 Arg Val Gln Phe Gln Ser Arg Asn Phe His Asn Ile Leu Gln Trp Gln
 35 40 45
 Pro Gly Arg Ala Leu Thr Gly Asn Ser Ser Val Tyr Phe Val Gln Tyr
 50 55 60
 Lys Ile Met Phe Ser Cys Ser Met Lys Ser Ser His Gln Lys Pro Ser
 65 70 75 80
 Gly Cys Trp Gln His Ile Ser Cys Asn Phe Pro Gly Cys Arg Thr Leu
 85 90 95
 Ala Lys Tyr Gly Gln Arg Gln Trp Lys Asn Lys Glu Asp Cys Trp Gly

			100					105				110					
Thr	Gln	Glu	Leu	Ser	Cys	Asp	Leu	Thr	Ser	Glu	Thr	Ser	Asp	Ile	Gln		
		115					120					125					
Glu	Pro	Tyr	Tyr	Gly	Arg	Val	Arg	Ala	Ala	Ser	Ala	Gly	Ser	Tyr	Ser		
	130					135					140						
Glu	Trp	Ser	Met	Thr	Pro	Arg	Phe	Thr	Pro	Trp	Trp	Glu	Thr	Lys	Ile		
145					150					155					160		
Asp	Pro	Pro	Val	Met	Asn	Ile	Thr	Gln	Val	Asn	Gly	Ser	Leu	Leu	Val		
			165					170					175				
Ile	Leu	His	Ala	Pro	Asn	Leu	Pro	Tyr	Arg	Tyr	Gln	Lys	Glu	Lys	Asn		
		180					185					190					
Val	Ser	Ile	Glu	Asp	Tyr	Tyr	Glu	Leu	Leu	Tyr	Arg	Val	Phe	Ile	Ile		
	195						200				205						
Asn	Asn	Ser	Leu	Glu	Lys	Glu	Gln	Lys	Val	Tyr	Glu	Gly	Ala	His	Arg		
	210					215				220							
Ala	Val	Glu	Ile	Glu	Ala	Leu	Thr	Pro	His	Ser	Ser	Tyr	Cys	Val	Val		
225					230					235					240		
Ala	Glu	Ile	Tyr	Gln	Pro	Met	Leu	Asp	Arg	Arg	Ser	Gln	Arg	Ser	Glu		
			245					250					255				
Glu	Arg	Cys	Val	Glu	Ile	Pro											
			260														